

Query Match                   99.6%; Score 1284; DB 6; Length 240;  
Best Local Similarity       99.6%; Pred. No. 2.9e-83;  
Matches 239; Conservative     0; Mismatches     1; Indels        0; Gaps        0;

Qy            1 MATAMYLEHYLDSIENLPCELQRNFQLMRELDQRTEDKKAÆIDILAAEYISTVKTLSPDQ 60  
              | | | | | | | | | | | | | | | | | | | | | | | | | | | |  
Db            1 MATAMYLEHYLDSIENLPCELQRNFQLMRELDQRTEDKKAÆIDILAAEYISTVKTLSPDQ 60

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Qy      61  RVERLQKIQNAYSCKEYSDDKVQLAMQTYEMVDKHIRRLDADLARFEADLKDKMEGSDF 120
        |||
Db      61  RVERLQKIQNAYSCKEYSDDKVQLAMQTYEMVDKHIRRLDADLARFEADLKDKMEGSDF 120

Qy     121  ESSGGRGLKKGRGQKEKRGSRGRGRRTSEEDTPK KKKKHKGSEFTDTILSVHPSDVLDMP 180
        |||
Db     121  ESSGGRGLKKGRGQKEKRGSRGRGRRTSEEDTPK KKKKHKGSEFTDTILPVHPSDVLDMP 180

Qy     181  VDPNEPTYCLCHQVSYGEMIGCDNPDCPIEWFHFACVDLTTPKGKWF CPRCVQEKRRKK 240
        |||
Db     181  VDPNEPTYCLCHQVSYGEMIGCDNPDCPIEWFHFACVDLTTPKGKWF CPRCVQEKRRKK 240
<!-- EndFragment -->

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